

FIG. 2A

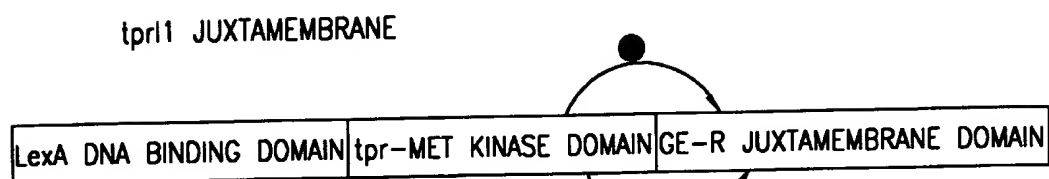


FIG. 2B

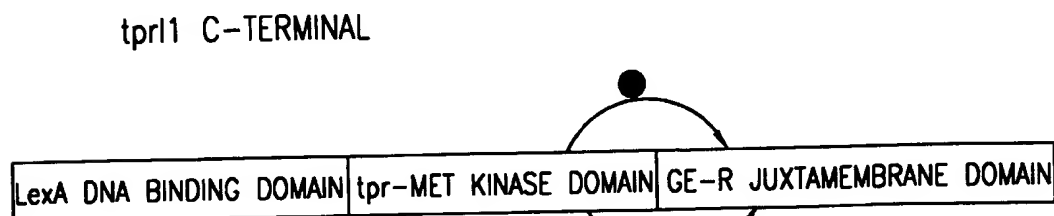


FIG. 2C

00017074-104504

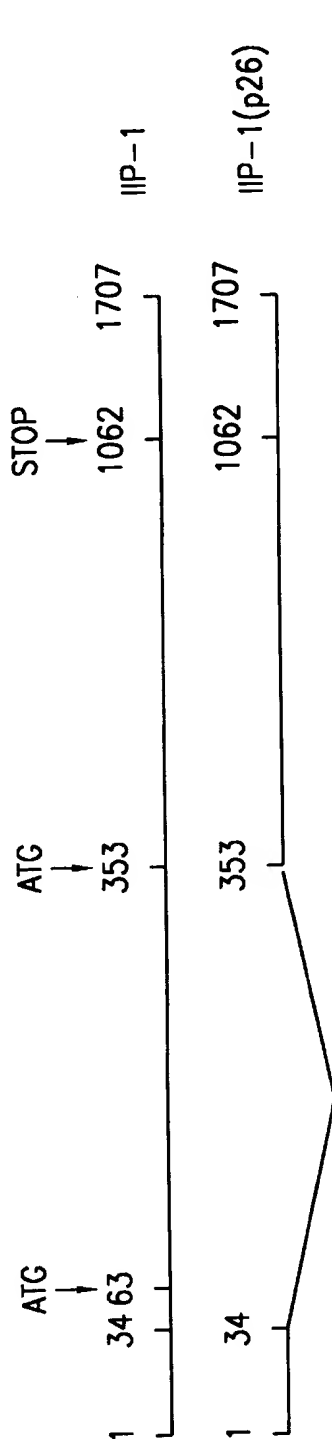


FIG. 3A

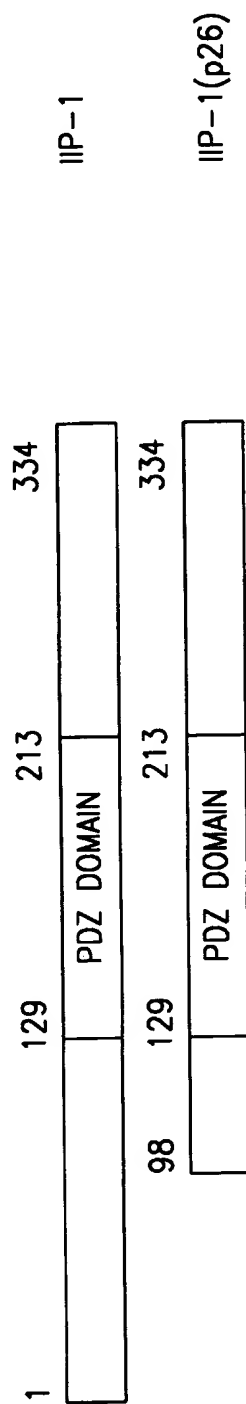


FIG. 3B

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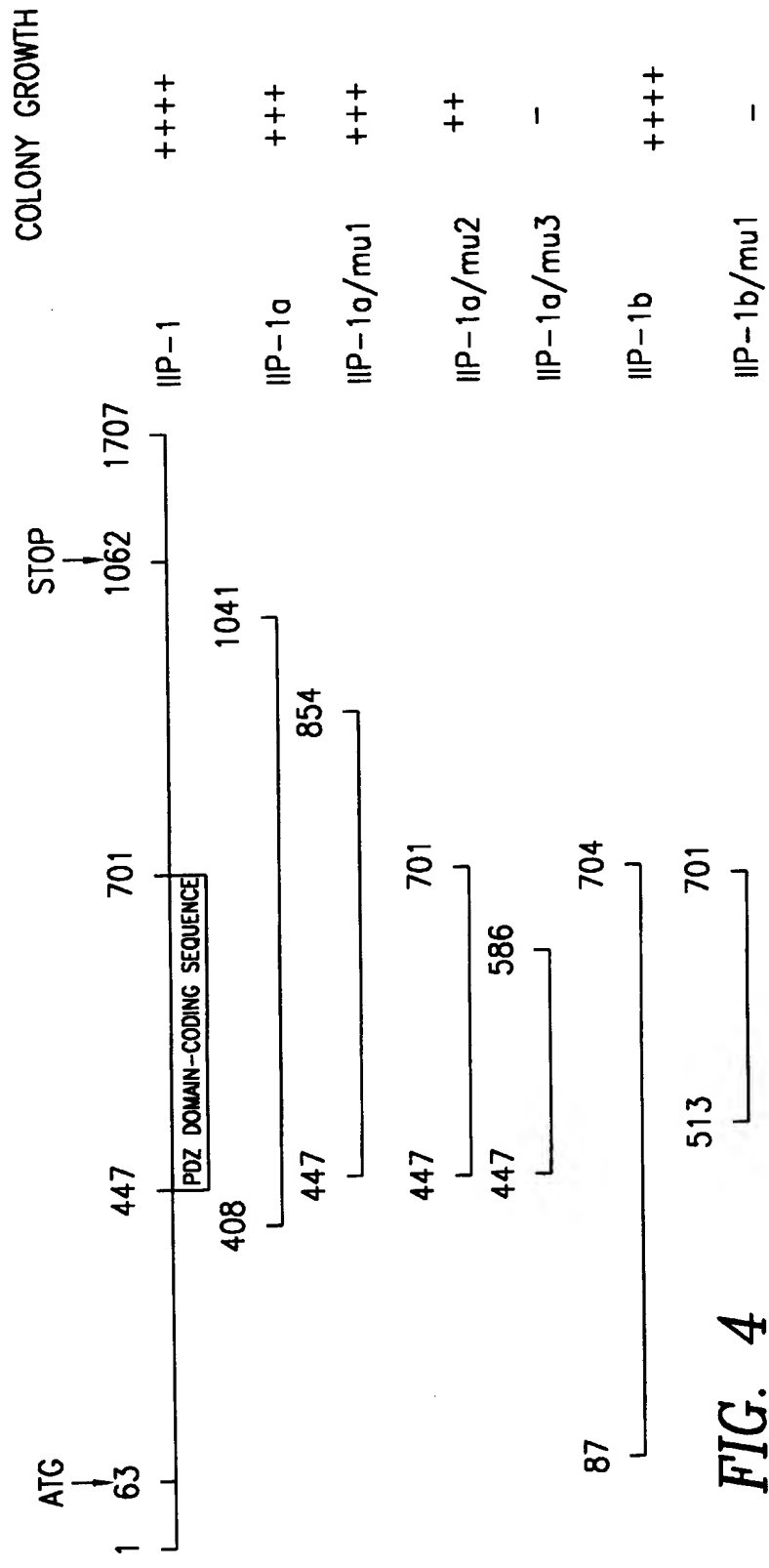


FIG. 4



Fig. 5

1 MSRPRLAG TSGDKGLSG KRTK<sup>+</sup>ENSEGE ALAKVEDSNP QKTSATKNCL

51 KNLSSHWMK SEPE<sup>+</sup>SRLEKG VDVKFSIEDL KAQPKQTTCW DGVRNYQARN

101 FLRAMKL<sup><</sup>GEE AFFYHSNCKE PGIAGLMKIV KEAYPDHTQF EKN<sup>+</sup>NP<sup>+</sup>HYDPS

151 SKEDNPKW<sup>+</sup>SM VDVQFVRMMK RFIPLAELKS YHQAHKATGG PLKNMVL<sup>+</sup>FTR

201 QRLSIQPLTQ EEFD<sup>+</sup>FLVLSLE EKEPS\*

- ▼ putative N-glycosylation site
- ◆ putative Casein kinase II phosphorylation site
- putative N-myristoylation site
- < putative Tyrosine kinase phosphorylation site
- + putative Protein kinase C phosphorylation site
- ↔ putative NLS